## SEQUENCE LISTING

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<110> Yan, Riqiang
      Tomasselli, Alfredo G.
      Gurney, Mark E.
      Emmons, Thomas L.
      Bienkowski, Mike J.
      Heinrikson, Robert L.
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Glu Glu Pro Glu Glu Pro Gly Arg Gly Ser Phe Val Glu Met Val
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Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr 65 70 75 80

Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser 85 90 95

Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
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Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp 130 135 140

Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile 145 150 155 160

Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp 165 170 175

Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro Asp Asp 180 185 190

Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His Val Pro 195 200 205

Asn Leu Phe Ser Leu His Leu Cys Gly Ala Gly Phe Pro Leu Asn Gln 210 215 220

Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly Ile 225 230 235 240

Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg Arg 245 250 255

Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly Gln 260 265 270

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Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
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Pro Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser 275 280 285

Ser Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val 290 295 300

Cys Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser 305 310 315 320

Leu Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile 325 330 335

Leu Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln 340 345 350

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Met Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala 370 380

Arg Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu 385 390 395 400

Phe Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu 405 410 415

Asp Cys Gly Tyr Asn Ile Pro Gln Thr Asp Glu Ser Thr Leu Met Thr 420 425 430

Ile Ala Tyr Val Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu 435 440 445

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peptide sequence

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  1
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Xaa Phe Ala Xaa Xaa Xaa Asn
<210> 111
<211> 8
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<220>
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      peptide sequence
<220>
<221> SITE
<222> (1)
<223> Xaa= any amino acid
<220>
<221> SITE
<222> (4)..(6)
<223> Xaa= any amino acid
<220>
<221> SITE
 <222> (7)
<223> Xaa= E, G, F, H, cysteic acid or S
<400> 111
 Xaa Phe Ala Xaa Xaa Xaa Asn
 1
 <210> 112
 <211> 8
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
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 <222> (1)
 <223> Xaa= any amino acid
 <220>
 <221> SITE
  <222> (4)..(7)
 <223> Xaa= any amino acid
  <220>
  <221> SITE
  <222> (8)
  <223> Xaa= F, W, G, A, H, P, G, N or S
  <400> 112
  Xaa Phe Ala Xaa Xaa Xaa Xaa Xaa
    1
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<210> 113
<211> 9
<212> PRT
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      peptide sequence
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Glu Val Asn Leu Asp Ala Glu Phe Arg
<210> 114
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 114
Asp Tyr Lys Asp Asp Asp Lys
<210> 115
<211> 17
<212> PRT
<213> Artificial Sequence
<220>
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      peptide sequence
 <400> 115
Ala Cys Gly Ser Glu Ser Met Asp Ser Gly Ile Ser Leu Asp Asn Lys
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                   5
 Trp
 <210> 116
 <211> 17
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 116
 Trp Lys Lys Gly Ala Ile Ile Gly Leu Met Val Gly Gly Val Val Lys
                                       10
 Lys
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<210> 117

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<211> 11
<212> PRT
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      peptide sequence
<400> 117
Ala Asn Leu Ser Thr Phe Ala Gln Pro Arg Arg
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<210> 118
<211> 22
<212> PRT
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      peptide sequence
<400> 118
Tyr Arg Tyr Gln Ser His Asp Tyr Ala Phe Ser Ser Val Glu Lys Leu
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Leu His Leu Gly Gly Cys
             20
<210> 119
<211> 22
<212> PRT
<213> Artificial Sequence
<220>
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      peptide sequence
<400> 119
Tyr Arg Tyr Gln Ser His Asp Tyr Ala Phe Ser Ser Val Glu Lys Leu
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 Leu His Leu Gly Gly Cys
 <210> 120
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 <212> PRT
 <213> Artificial Sequence
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 Lys Thr Ile Thr Leu Glu Val Glu Pro Ser
                   5
 <210> 121
 <211> 12
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<220>
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<222> (9)
<223> Xaa= cysteic acid
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<210> 122
<211> 11
<212> PRT
<213> Artificial Sequence
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      peptide sequence
<400> 122
Val Glu Ala Leu Tyr Leu Val Glu Gly Glu Arg
<210> 123
<211> 363
<212> PRT
<213> Homo sapiens
 <220>
 <223> galactosyltransferase
 <400> 123
 Met Ala Ser Lys Ser Trp Leu Asn Phe Leu Thr Phe Leu Cys Gly Ser
 Ala Ile Gly Phe Leu Leu Cys Ser Gln Leu Phe Ser Ile Leu Leu Gly
 Glu Lys Val Asp Thr Gln Pro Asn Val Leu His Asn Asp Pro His Ala
 Arg His Ser Asp Asp Asn Gly Gln Asn His Leu Glu Gly Gln Met Asn
 Phe Asn Ala Asp Ser Ser Gln His Lys Asp Glu Asn Thr Asp Ile Ala
 Glu Asn Leu Tyr Gln Lys Val Arg Ile Leu Cys Trp Val Met Thr Gly
 Pro Gln Asn Leu Glu Lys Lys Ala Lys His Val Lys Ala Thr Trp Ala
                                  105
 Gln Arg Cys Asn Lys Val Leu Phe Met Ser Ser Glu Glu Asn Lys Asp
```

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Phe Pro Ala Val Gly Leu Lys Thr Lys Glu Gly Arg Asp Gln Leu Tyr Trp Lys Thr Ile Lys Ala Phe Gln Tyr Val His Glu His Tyr Leu Glu Asp Ala Asp Trp Phe Leu Lys Ala Asp Asp Asp Thr Tyr Val Ile Leu Asp Asn Leu Arg Trp Leu Leu Ser Lys Tyr Asp Pro Glu Glu Pro Ile Tyr Phe Gly Arg Arg Phe Lys Pro Tyr Val Lys Gln Gly Tyr Met Ser Gly Gly Ala Gly Tyr Val Leu Ser Lys Glu Ala Leu Lys Arg Phe Val Asp Ala Phe Lys Thr Asp Lys Cys Thr His Ser Ser Ser Ile Glu Asp 230 Leu Ala Leu Gly Arg Cys Met Glu Ile Met Asn Val Glu Ala Gly Asp 250 Ser Arg Asp Thr Ile Gly Lys Glu Thr Phe His Pro Phe Val Pro Glu 265 His His Leu Ile Lys Gly Tyr Leu Pro Arg Thr Phe Trp Tyr Trp Asn Tyr Asn Tyr Tyr Pro Pro Val Glu Gly Pro Gly Cys Cys Ser Asp Leu 300 Ala Val Ser Phe His Tyr Val Asp Ser Thr Thr Met Tyr Glu Leu Glu 305 310 Tyr Leu Val Tyr His Leu Arg Pro Tyr Gly Tyr Leu Tyr Arg Tyr Gln 330 Pro Thr Leu Pro Glu Arg Ile Leu Lys Glu Ile Ser Gln Ala Asn Lys Asn Glu Asp Thr Lys Val Lys Leu Gly Asn Pro <210> 124

<211> 405

<212> PRT

<213> Homo sapiens

<220>

<223> Homo sapiens sialylytransferase 1

<400> 124

Ile His Thr Asn Leu Lys Lys Lys Phe Ser Cys Cys Val Leu Val Phe

Leu Leu Phe Ala Val Ile Cys Val Trp Lys Glu Lys Lys Gly Ser

Tyr Tyr Asp Ser Phe Lys Leu Gln Thr Lys Glu Phe Gln Val Leu Lys

| Ser        | Leu<br>50  | Gly        | Lys        | Leu        | Ala        | Met<br>55  | Gly        | Ser        | Asp        | Ser        | Gln<br>60  | Ser        | Val        | Ser        | Ser        |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ser<br>65  | Ser        | Thr        | Gln        | Asp        | Pro<br>70  | His        | Arg        | Gly        | Arg        | Gln<br>75  | Thr        | Leu        | Gly        | Ser        | Leu<br>80  |
| Arg        | Gly        | Leu        | Ala        | Lys<br>85  | Ala        | Lys        | Pro        | Glu        | Ala<br>90  | Ser        | Phe        | Gln        | Val        | Trp<br>95  | Asn        |
| Lys        | Asp        | Ser        | Ser<br>100 | Ser        | Lys        | Asn        | Leu        | Ile<br>105 | Pro        | Arg        | Leu        | Gln        | Lys<br>110 | Ile        | Trp        |
| Lys        | Asn        | Tyr<br>115 | Leu        | Ser        | Met        | Asn        | Lys<br>120 | Tyr        | Lys        | Val        | Ser        | Tyr<br>125 | Lys        | Gly        | Pro        |
| Gly        | Pro<br>130 | Gly        | Ile        | Lys        | Phe        | Ser<br>135 | Ala        | Glu        | Ala        | Leu        | Arg<br>140 | Cys        | His        | Leu        | Arg        |
| Asp<br>145 | His        | Val        | Asn        | Val        | Ser<br>150 | Met        | Val        | Glu        | Val        | Thr<br>155 | Asp        | Phe        | Pro        | Phe        | Asn<br>160 |
| Thr        | Ser        | Glu        | Trp        | Glu<br>165 | Gly        | Tyr        | Leu        | Pro        | Lys<br>170 | Glu        | Ser        | Ile        | Arg        | Thr<br>175 | Lys        |
| Ala        | Gly        | Pro        | Trp<br>180 | Gly        | Arg        | Cys        | Ala        | Val<br>185 | Val        | Ser        | Ser        | Ala        | Gly<br>190 | Ser        | Leu        |
| Lys        | Ser        | Ser<br>195 | Gln        | Leu        | Gly        | Arg        | Glu<br>200 | Ile        | Asp        | Asp        | His        | Asp<br>205 | Ala        | Val        | Leu        |
| Arg        | Phe<br>210 | Asn        | Gly        | Ala        | Pro        | Thr<br>215 | Ala        | Asn        | Phe        | Gln        | Gln<br>220 | Asp        | Val        | Gly        | Thr        |
| Lys<br>225 | Thr        | Thr        | Ile        | Arg        | Leu<br>230 | Met        | Asn        | Ser        | Gln        | Leu<br>235 | Val        | Thr        | Thr        | Glu        | Lys<br>240 |
| Arg        | Phe        | Leu        | Lys        | Asp<br>245 | Ser        | Leu        | Tyr        | Asn        | Glu<br>250 | Gly        | Ile        | Leu        | Ile        | Val<br>255 | Trp        |
| Asp        | Pro        | Ser        | Val<br>260 |            | His        | Ser        | Asp        | Ile<br>265 |            | Lys        | Trp        | Tyr        | Gln<br>270 | Asn        | Pro        |
| Asp        | Tyr        | Asn<br>275 |            | Phe        | Asn        | Asn        | Tyr<br>280 |            | Thr        | Tyr        | Arg        | Lys<br>285 | Leu        | His        | Pro        |
| Asn        | Gln<br>290 |            | Phe        | Tyr        | Ile        | Leu<br>295 |            | Pro        | Gln        | . Met      | Pro<br>300 |            | Glu        | Leu        | Trp        |
| Asp<br>305 |            | Leu        | Gln        | Glu        | Ile<br>310 |            | Pro        | Glu        | Glu        | 1le<br>315 | Gln        | Pro        | Asn        | Pro        | Pro<br>320 |
| Ser        | Ser        | Gly        | Met        | Leu<br>325 |            | Ile        | lle        | Ile        | Met<br>330 |            | Thr        | Leu        | . Cys      | Asp<br>335 | Gln        |
| Val        | Asp        | Ile        | Tyr<br>340 |            | Phe        | Leu        | Pro        | Ser<br>345 |            | Arg        | Lys        | Thr        | Asp<br>350 | Val        | Cys        |
| Туг        | Туг        | Туг<br>355 |            | Lys        | s Ph∈      | Ph∈        | 360        |            | Ala        | Cys        | Thr        | Met<br>365 | Gly        | Ala        | Tyr        |
| His        | Pro        | Lev        | . Lev      | ι Туг      | Glu        | Lys        | . Asn      | Lev        | ı Val      | Lys        | His        | Leu        | Asn        | Gln        | Gly        |

Thr Asp Glu Asp Ile Tyr Leu Leu Gly Lys Ala Thr Leu Pro Gly Phe 385 390 395

Arg Thr Ile His Cys 405

<210> 125

<211> 518

<212> PRT

<213> Homo sapiens

<220>

<223> Homo sapiens aspartyl protease 1

<400> 125

Met Gly Ala Leu Ala Arg Ala Leu Leu Leu Pro Leu Leu Ala Gln Trp

1 5 10 15

Leu Leu Arg Ala Ala Pro Glu Leu Ala Pro Ala Pro Phe Thr Leu Pro 20 25 30

Leu Arg Val Ala Ala Ala Thr Asn Arg Val Val Ala Pro Thr Pro Gly

Pro Gly Thr Pro Ala Glu Arg His Ala Asp Gly Leu Ala Leu 50 55 60

Glu Pro Ala Leu Ala Ser Pro Ala Gly Ala Ala Asn Phe Leu Ala Met
65 70 75 80

Val Asp Asn Leu Gln Gly Asp Ser Gly Arg Gly Tyr Tyr Leu Glu Met 85 90 95

Leu Ile Gly Thr Pro Pro Gln Lys Leu Gln Ile Leu Val Asp Thr Gly 100 105 110

Ser Ser Asn Phe Ala Val Ala Gly Thr Pro His Ser Tyr Ile Asp Thr 115 120 125

Tyr Phe Asp Thr Glu Arg Ser Ser Thr Tyr Arg Ser Lys Gly Phe Asp 130 135 140

Val Thr Val Lys Tyr Thr Gln Gly Ser Trp Thr Gly Phe Val Gly Glu 145 150 155 160

Asp Leu Val Thr Ile Pro Lys Gly Phe Asn Thr Ser Phe Leu Val Asn 165 170 175

Ile Ala Thr Ile Phe Glu Ser Glu Asn Phe Phe Leu Pro Gly Ile Lys
180 185 190

Trp Asn Gly Ile Leu Gly Leu Ala Tyr Ala Thr Leu Ala Lys Pro Ser

Ser Ser Leu Glu Thr Phe Phe Asp Ser Leu Val Thr Gln Ala Asn Ile 210 215 220

Pro Asn Val Phe Ser Met Gln Met Cys Gly Ala Gly Leu Pro Val Ala 225 230 235 240 Gly Ser Gly Thr Asn Gly Gly Ser Leu Val Leu Gly Gly Ile Glu Pro  $245 \hspace{1cm} 250 \hspace{1cm} 255$ 

Ser Leu Tyr Lys Gly Asp Ile Trp Tyr Thr Pro Ile Lys Glu Glu Trp 260 265 270

Tyr Tyr Gln Ile Glu Ile Leu Lys Leu Glu Ile Gly Gly Gln Ser Leu 275 280 285

Asn Leu Asp Cys Arg Glu Tyr Asn Ala Asp Lys Ala Ile Val Asp Ser 290 295 300

Gly Thr Thr Leu Leu Arg Leu Pro Gln Lys Val Phe Asp Ala Val Val 305 310 315

Glu Ala Val Ala Arg Ala Ser Leu Ile Pro Glu Phe Ser Asp Gly Phe 325 330 335

Trp Thr Gly Ser Gln Leu Ala Cys Trp Thr Asn Ser Glu Thr Pro Trp 340 345 350

Ser Tyr Phe Pro Lys Ile Ser Ile Tyr Leu Arg Asp Glu Asn Ser Ser 355 360 365

Arg Ser Phe Arg Ile Thr Ile Leu Pro Gln Leu Tyr Ile Gln Pro Met 370 380

Met Gly Ala Gly Leu Asn Tyr Glu Cys Tyr Arg Phe Gly Ile Ser Pro 385 390 395 400

Ser Thr Asn Ala Leu Val Ile Gly Ala Thr Val Met Glu Gly Phe Tyr 405 410 415

Val Ile Phe Asp Arg Ala Gln Lys Arg Val Gly Phe Ala Ala Ser Pro 420 425 430

Cys Ala Glu Ile Ala Gly Ala Ala Val Ser Glu Ile Ser Gly Pro Phe 435 440 445

Ser Thr Glu Asp Val Ala Ser Asn Cys Val Pro Ala Gln Ser Leu Ser 450 455 460

Glu Pro Ile Leu Trp Ile Val Ser Tyr Ala Leu Met Ser Val Cys Gly
465 470 475 480

Ala Ile Leu Leu Val Leu Ile Val Leu Leu Leu Leu Pro Phe Arg Cys
485
490
495

Gln Arg Arg Pro Arg Asp Pro Glu Val Val Asn Asp Glu Ser Ser Leu 500 505 510

Val Arg His Arg Trp Lys 515

<210> 126

<211> 255

<212> PRT

<213> Homo sapiens

<220>

<223> Homo sapiens syntaxin 6

<400> 126 Met Ser Met Glu Asp Pro Phe Phe Val Val Lys Gly Glu Val Gln Lys Ala Val Asn Thr Ala Gln Gly Leu Phe Gln Arg Trp Thr Glu Leu Leu Gln Asp Pro Ser Thr Ala Thr Arg Glu Glu Ile Asp Trp Thr Thr Asn Glu Leu Arg Asn Asn Leu Arg Ser Ile Glu Trp Asp Leu Glu Asp Leu Asp Glu Thr Ile Ser Ile Val Glu Ala Asn Pro Arg Lys Phe Asn Leu Asp Ala Thr Glu Leu Ser Ile Arg Lys Ala Phe Ile Thr Ser Thr Arg Gln Val Val Arg Asp Met Lys Asp Gln Met Ser Thr Ser Ser Val Gln Ala Leu Ala Glu Arg Lys Asn Arg Gln Ala Leu Leu Gly Asp Ser Gly Ser Gln Asn Trp Ser Thr Gly Thr Thr Asp Lys Tyr Gly Arg Leu Asp 135 Arg Glu Leu Gln Arg Ala Asn Ser His Phe Ile Glu Glu Gln Gln Ala 150 Gln Gln Gln Leu Ile Val Glu Gln Gln Asp Glu Gln Leu Glu Leu Val 170 Ser Gly Ser Ile Gly Val Leu Lys Asn Met Ser Gln Arg Ile Gly Gly Glu Leu Glu Glu Gln Ala Val Met Leu Glu Asp Phe Ser His Glu Leu 200 Glu Ser Thr Gln Ser Arg Leu Asp Asn Val Met Lys Lys Leu Ala Lys 210 Val Ser His Met Thr Ser Asp Arg Arg Gln Trp Cys Ala Ile Ala Ile

<210> 127 <211> 1728

<212> DNA <213> Artificial Sequence

245

<220>

<223> Description of Artificial Sequence: nucleic acid encoding recombinant fusion protein

Leu Phe Ala Val Leu Leu Val Val Leu Ile Leu Phe Leu Val Leu

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gttgaggagg agaacccgga cttctggaac cgcgaggcag ccgaggccct gggtgccgcc 120
aagaagctgc agcctgcaca gacagccgcc aagaacctca tcatcttcct gggcgatggg 180

atgggggtgt ctacggtgac agctgccagg atcctaaaag ggcagaagaa ggacaaactg 240

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gggcctgaga tacccctggc catggaccgc ttcccatatg tggctctgtc caagacatac 300
aatgtagaca aacatgtgcc agacagtgga gccacagcca cggcctacct gtgcggggtc 360
aagggcaact tecagaceat tggettgagt geageegeee getttaacea gtgcaacaeg 420
acacgoggca acgaggtcat ctccgtgatg aatcgggcca agaaagcagg gaagtcagtg 480
ggagtggtaa ccaccacacg agtgcagcac gcctcgccag ccggcaccta cgcccacacg 540
gtgaaccgca actggtactc ggacgccgac gtgcctgcct cggcccgcca ggaggggtgc 600
caggacateg ctacgcaget catetecaac atggacattg acgtgateet aggtggagge 660
cgaaagtaca tgtttcccat gggaacccca gaccctgagt acccagatga ctacagccaa 720
ggtgggacca ggctggacgg gaagaatctg gtgcaggaat ggctggcgaa gcgccagggt 780
geceggtatg tgtggaaceg cactgagete atgeaggett ceetggacee gtetgtgace 840
catctcatgg gtctctttga gcctggagac atgaaatacg agatccaccg agactccaca 900
ctggacccct ccctgatgga gatgacagag gctgccctgc gcctgctgag caggaacccc 960
cgcggcttct tcctcttcgt ggagggtggt cgcatcgacc atggtcatca tgaaagcagg 1020
gettaceggg caetgaetga gaegateatg ttegaegaeg ceattgagag ggegggeeag 1080
ctcaccageg aggaggacae getgageete gtcactgeeg accaeteeca egtettetee 1140
tteggagget acceettgeg agggagetee atetteggge tggeeeetgg caaggeeegg 1200
gacaggaagg cctacacggt cctcctatac ggaaacggtc caggctatgt gctcaaggac 1260
ggcgcccggc cggatgttac cgagagcgag agcgggagcc ccgagtatcg gcagcagtca 1320
gcagtgcccc tggacgaaga gacccacgca ggcgaggacg tggcggtgtt cgcgcgcggc 1380
cegcaggege acetggttea eggegtgeag gageagacet teatagegea egteatggee 1440
ttcgccgcct gcctggagcc ctacaccgcc tgcgacctgg cgccccccgc cggcaccacc 1500
gacgccgcgc acccaggtaa ctatgaagtt gaattccgaa gagcactcta cgtagagggt 1560
gaaagaggat tettetacae tecaaaggea etetaceteg tagagggtga aagaggatte 1620
ttctacacta gtctcatgac catagcctat gtcatggctg ccatctgcgc cctcttcatg 1680
ctgccactct gcctcatggt ggactacaag gatgatgatg acaagtag
<210> 128
<211> 575
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: recombinant
       fusion protein sequence
 <400> 128
Met Leu Leu Leu Leu Leu Gly Leu Arg Leu Gln Leu Ser Leu
 Gly Ile Ile Pro Val Glu Glu Glu Asn Pro Asp Phe Trp Asn Arg Glu
 Ala Ala Glu Ala Leu Gly Ala Ala Lys Lys Leu Gln Pro Ala Gln Thr
 Ala Ala Lys Asn Leu Ile Ile Phe Leu Gly Asp Gly Met Gly Val Ser
 Thr Val Thr Ala Ala Arg Ile Leu Lys Gly Gln Lys Lys Asp Lys Leu
 Gly Pro Glu Ile Pro Leu Ala Met Asp Arg Phe Pro Tyr Val Ala Leu
 Ser Lys Thr Tyr Asn Val Asp Lys His Val Pro Asp Ser Gly Ala Thr
                                                     110
                                 105
             100
 Ala Thr Ala Tyr Leu Cys Gly Val Lys Gly Asn Phe Gln Thr Ile Gly
                             120
 Leu Ser Ala Ala Ala Arg Phe Asn Gln Cys Asn Thr Thr Arg Gly Asn
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135

| Glu<br>145 | Val        | Ile        | Ser        | Val        | Met<br>150 | Asn        | Arg        | Ala        | Lys        | Lys<br>155 | Ala        | Gly        | Lys        | Ser        | Val<br>160 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gly        | Val        | Val        | Thr        | Thr<br>165 | Thr        | Arg        | Val        | Gln        | His<br>170 | Ala        | Ser        | Pro        | Ala        | Gly<br>175 | Thr        |
| Tyr        | Ala        | His        | Thr<br>180 | Val        | Asn        | Arg        | Asn        | Trp<br>185 | Tyr        | Ser        | Asp        | Ala        | Asp<br>190 | Val        | Pro        |
| Ala        | Ser        | Ala<br>195 | Arg        | Gln        | Glu        | Gly        | Cys<br>200 | Gln        | Asp        | Ile        | Ala        | Thr<br>205 | Gln        | Leu        | Ile        |
| Ser        | Asn<br>210 | Met        | Asp        | Ile        | Asp        | Val<br>215 | Ile        | Leu        | Gly        | Gly        | Gly<br>220 | Arg        | Lys        | Tyr        | Met        |
| Phe<br>225 | Pro        | Met        | Gly        | Thr        | Pro<br>230 | Asp        | Pro        | Glu        | Tyr        | Pro<br>235 | Asp        | Asp        | Tyr        | Ser        | Gln<br>240 |
| Gly        | Gly        | Thr        | Arg        | Leu<br>245 | Asp        | Gly        | Lys        | Asn        | Leu<br>250 | Val        | Gln        | Glu        | Trp        | Leu<br>255 | Ala        |
| Lys        | Arg        | Gln        | Gly<br>260 | Ala        | Arg        | Tyr        | Val        | Trp<br>265 | Asn        | Arg        | Thr        | Glu        | Leu<br>270 | Met        | Gln        |
| Ala        | Ser        | Leu<br>275 | Asp        | Pro        | Ser        | Val        | Thr<br>280 | His        | Leu        | Met        | Gly        | Leu<br>285 | Phe        | Glu        | Pro        |
| _          | 290        |            | _          |            |            | 295        |            |            |            |            | 300        |            |            | Pro        |            |
| 305        |            |            |            |            | 310        |            |            |            |            | 315        |            |            |            | Asn        | 320        |
|            |            |            |            | 325        |            |            |            |            | 330        |            |            |            |            | Gly<br>335 |            |
| His        | Glu        | Ser        | Arg<br>340 | Ala        | Tyr        | Arg        | Ala        | Leu<br>345 | Thr        | Glu        | Thr        | Ile        | Met<br>350 | Phe        | Asp        |
| Asp        | Ala        | Ile<br>355 | Glu        | Arg        | Ala        | Gly        | Gln<br>360 | Leu        | Thr        | Ser        | Glu        | Glu<br>365 | Asp        | Thr        | Leu        |
| Ser        | Leu<br>370 | Val        | Thr        | Ala        | Asp        | His<br>375 | Ser        | His        | Val        | Phe        | Ser<br>380 | Phe        | Gly        | Gly        | Tyr        |
| Pro<br>385 | Leu        | Arg        | Gly        | Ser        | Ser<br>390 | Ile        | Phe        | Gly        | Leu        | Ala<br>395 | Pro        | Gly        | Lys        | Ala        | Arg<br>400 |
| Asp        | Arg        | Lys        | Ala        | Tyr<br>405 | Thr        | Val        | Leu        | Leu        | Tyr<br>410 | Gly        | Asn        | Gly        | Pro        | Gly<br>415 | Tyr        |
| Val        | Leu        | Lys        | Asp<br>420 | Gly        | Ala        | Arg        | Pro        | Asp<br>425 | Val        | Thr        | Glu        | Ser        | Glu<br>430 | Ser        | Gly        |
| Ser        | Pro        | Glu<br>435 | Tyr        | Arg        | Gln        | Gln        | Ser<br>440 | Ala        | Val        | Pro        | Leu        | Asp<br>445 | Glu        | Glu        | Thr        |
| His        | Ala<br>450 | Gly        | Glu        | Asp        | Val        | Ala<br>455 | Val        | Phe        | Ala        | Arg        | Gly<br>460 | Pro        | Gln        | Ala        | His        |
| Leu<br>465 | Val        | His        | Gly        | Val        | Gln<br>470 | Glu        | Gln        | Thr        | Phe        | Ile<br>475 | Ala        | His        | Val        | Met        | Ala<br>480 |

Phe Ala Ala Cys Leu Glu Pro Tyr Thr Ala Cys Asp Leu Ala Pro Pro 490 Ala Gly Thr Thr Asp Ala Ala His Pro Gly Asn Tyr Glu Val Glu Pro Arg Arg Ala Leu Tyr Val Glu Gly Glu Arg Gly Phe Phe Tyr Thr Pro 520 Lys Ala Leu Tyr Leu Val Glu Gly Glu Arg Gly Phe Phe Tyr Thr Ser 530 Leu Met Thr Ile Ala Tyr Val Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Asp Tyr Lys Asp Asp Asp Asp Lys <210> 129 <211> 5 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: synthetic peptide sequence <400> 129 Lys Met Asp Ala Glu <210> 130 <211> 5 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic peptide sequence <400> 130 Gly Arg Arg Gly Ser <210> 131 <211> 10 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: synthetic peptide sequence <400> 131 Val Glu Ala Asn Tyr Glu Val Glu Gly Glu 5

<210> 132 <211> 10

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<212> PRT
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      peptide sequence
 <400> 132
Val Glu Ala Asn Tyr Ala Val Glu Gly Glu
                   5
<210> 133
<211> 10
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<213> Artificial Sequence
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      peptide sequence
<400> 133
Lys Thr Ile Asn Leu Glu Val Glu Pro Ser
 1
                  5
<210> 134
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
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<222> (5)
<223> Nle
<400> 134
Lys Thr Ile Asn Xaa Glu Val Glu Pro Ser
<210> 135
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<221> MOD RES
<222> (5)
<223> Nle
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 135
Lys Thr Ile Asn Xaa Glu Val Asp Pro Ser
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<210> 136
<211> 10
<212> PRT
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<220>
<221> MOD_RES
<222> (5)
<223> Nle
<220>
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      peptide sequence
<400> 136
Lys Thr Ile Asn Xaa Asp Val Asp Pro Ser
<210> 137
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<400> 137
Lys Thr Ile Ser Leu Asp Val Glu Pro Ser
                   5
 1
<210> 138
<211> 10
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 <213> Artificial Sequence
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       peptide sequence
 <400> 138
 Lys Thr Ile Ser Leu Asp Val Asp Pro Ser
                   5
   1
 <210> 139
 <211> 4
 <212> PRT
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       peptide sequence
 <400> 139
 Lys Met Asp Ala
   1
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<210> 140
<211> 4
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 140
Ser Tyr Glu Val
<210> 141
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 141
 Ser Glu Val Ser Tyr Glu Val Glu Phe Arg
 1
 <210> 142
 <211> 4
 <212> PRT
 <213> Artificial Sequence
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 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 142
 Asn Leu Asp Ala
   1
 <210> 143
 <211> 10
  <212> PRT
  <213> Artificial Sequence
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  <220>
        peptide sequence
  <400> 143
  Ser Glu Val Ser Tyr Asp Ala Glu Phe Arg
  <210> 144
  <211> 10
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: synthetic
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peptide sequence

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<400> 144
Ser Glu Val Ser Tyr Glu Ala Glu Phe Arg
<210> 145
<211> 25
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
<220>
      peptide sequence
Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser
 Glu Val Ser Tyr Glu Val Glu Phe Arg
              20
 <210> 146
 <211> 20
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: synthetic
 <220>
       peptide sequence
 Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Val Ser Tyr Glu
 Val Glu Phe Arg
  <210> 147
  <211> 15
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
  Lys Thr Glu Glu Ile Ser Glu Val Ser Tyr Glu Val Glu Phe Arg
                                        10
                    5
    1
  <210> 148
  <211> 10
  <212> PRT
  <213> Artificial Sequence
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  <223> Description of Artificial Sequence: synthetic
        peptide sequence
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<400> 148
Thr Glu Val Ser Tyr Glu Val Glu Phe Arg
<210> 149
<211> 10
<212> PRT
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      peptide sequence
<400> 149
Ser Glu Val Asp Tyr Glu Val Glu Phe Arg
 <210> 150
 <211> 10
 <212> PRT
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 150
 Thr Glu Val Asp Tyr Glu Val Glu Phe Arg
 <210> 151
 <211> 10
 <212> PRT
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  <223> Description of Artificial Sequence: synthetic
        peptide sequence
  <400> 151
  Thr Glu Ile Asp Tyr Glu Val Glu Phe Arg
    1
  <210> 152
  <211> 10
  <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
   <400> 152
   Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg
                     5
   <210> 153
   <211> 10
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<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
Ser Glu Ile Asp Tyr Glu Val Glu Phe Arg
<210> 154
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (11)
<223> Xaa=tryptophan
<220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 154
 Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
   1
 <210> 155
 <211> 18
 <212> PRT
 <213> Artificial Sequence
 <220>
 <221> SITE
 <222> (16)
 <223> Xaa=tryptophan
 <223> Description of Artificial Sequence: synthetic
 <220>
       peptide sequence
  Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Kaa
  Lys Lys
  <210> 156
  <211> 23
  <212> PRT
  <213> Artificial Sequence
  <220>
  <221> SITE
  <222> (21)
  <223> Xaa=tryptophan
  <220>
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<223> Description of Artificial Sequence: synthetic
     peptide sequence
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val
                                      10
Glu Phe Arg Xaa Lys Lys
        20
<210> 157
<211> 28
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<220>
<221> SITE
<222> (26)
<223> Xaa=tryptophan
 <400> 157
Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser
 Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
 <210> 158
 <211> 13
 <212> PRT
 <213> Artificial Sequence
 <220>
 <221> SITE
 <222> (11)
 <223> Xaa=tryptophan
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
  <210> 159
  <211> 18
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
  <220>
  <221> SITE
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<222> (16)
<223> Xaa=tryptophan
Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg
  1
Xaa Lys Lys
<210> 160
<211> 23
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (21)
<223> Xaa=tryptophan
<223> Description of Artificial Sequence: synthetic
      peptide
 Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr
 Glu Val Glu Phe Arg Xaa Lys Lys
 <210> 161
 <211> 28
 <212> PRT
 <213> Artificial Sequence
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  <222> (26)
 <223> Xaa=tryptophan
  <220>
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
  Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile
  <400> 161
                                        10
  Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
  <210> 162
  <211> 13
  <212> PRT
  <213> Artificial Sequence
  <220>
  <221> SITE
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<222> (11)
<223> Xaa=oregon green
<223> Description of Artificial Sequence: synthetic
      peptide sequence
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
<210> 163
<211> 18
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
 <222> (16)
 <223> Xaa=oregon green
 <220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Kaa
                   5
  1
 Lys Lys
 <210> 164
 <211> 23
 <212> PRT
 <213> Artificial Sequence
 <220>
 <221> SITE
 <222> (21)
  <223> Xaa=oregon green
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
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  <400> 164
  Val Glu Phe Arg Xaa Lys Lys
               20
  <210> 165
  <211> 28
  <212> PRT
  <213> Artificial Sequence
   <220>
   <221> SITE
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<222> (26)
<223> Xaa=oregon green
<223> Description of Artificial Sequence: synthetic peptide sequence
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Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
<210> 166
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<221> SITE
<222> (11)
<223> Xaa=oregon green
<220>
<223> Description of Artificial Sequence: synthetic
       peptide sequence
 Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
 <210> 167
 <211> 18
 <212> PRT
 <213> Artificial Sequence
 <220>
 <221> SITE
 <222> (16)
 <223> Xaa=oregon green
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
  Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg
  <400> 167
                                        10
  Xaa Lys Lys
  <210> 168
  <211> 23
  <212> PRT
  <213> Artificial Sequence
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<220>

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<221> SITE
<222> (21)
<223> Xaa=oregon green
<223> Description of Artificial Sequence: synthetic
      peptide sequence
Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr
                  5
Glu Val Glu Phe Arg Xaa Lys Lys
                20
<210> 169
<211> 28
<212> PRT
<213> Artificial Sequence
 <220>
 <221> SITE
 <222> (26)
 <223> Xaa=oregon green
 <223> Description of Artificial Sequence: synthetic
 <220>
       peptide sequence
 Thr Arg Pro Gly Ser Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile
 Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Xaa Lys Lys
                  20
 <210> 170
 <211> 10
 <212> PRT
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
  <400> 170
  Ser Glu Val Asn Tyr Glu Val Glu Phe Arg
  <210> 171
  <211> 47
  <212> DNA
  <213> Artificial Sequence
  <220>
  <223> Description of Artificial Sequence: synthetic
        primer for site-directed mutagenesis of APP
   <400> 171
  gagatetetg aaattagtta tgaagtagaa tteegacatg aeteagg
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| <210> 172<br><211> 48<br><212> DNA<br><213> Artificial Sequence  |    |
|--|----|
| <220> <223> Description of Artificial Sequence: synthetic primer for site-directed mutagenesis of APP                            |    |
| <400> 172<br>tgagtcatgt cggaattcta cttcataact aatttcagag atctcctc  | 48 |
| <210> 173<br><211> 47<br><212> DNA<br><213> Artificial Sequence  |    |
| <pre>&lt;220&gt; &lt;223&gt; Description of Artificial Sequence: synthetic     primer for site-directed mutagenesis of APP</pre> |    |
| <400> 173 gagatetetg aaagtagtta tgaagtagaa tteegacatg aeteagg  | 47 |
| <210> 174<br><211> 48<br><212> DNA<br><213> Artificial Sequence  |    |
| <pre>&lt;220&gt; &lt;223&gt; Description of Artificial Sequence: synthetic     primer for site-directed mutagenesis of APP</pre> |    |
| <400> 174 tgagtcatgt cggaattcta cttcataact actttcagag atctcctc   | 48 |
| <210> 175<br><211> 47<br><212> DNA<br><213> Artificial Sequence  |    |
| <220> <223> Description of Artificial Sequence: synthetic primer for site-directed mutagenesis of APP                            |    |
| <400> 175 gagatetetg aaattagtta tgaageagaa tteegaeatg aeteagg  | 47 |
| <210> 176<br><211> 48<br><212> DNA<br><213> Artificial Sequence  |    |
| <220> <223> Description of Artificial Sequence: synthetic primer for site-directed mutagenesis of APP                            |    |
| <400> 176 tgagtcatgt cggaattetg etteataaet aattteagag ateteete   | 48 |
| <210> 177 <211> 5 <212> PRT  |    |

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<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
<400> 177
Val Ser Tyr Glu Val
<210> 178
<211> 5
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<213> Artificial Sequence
<220>
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 178
 Val Ser Tyr Asp Ala
  1
 <210> 179
 <211> 5
 <212> PRT
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 <220>
       peptide sequence
 <400> 179
 Ile Ser Tyr Glu Val
  <210> 180
  <211> 5
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
  <400> 180
  Val Lys Met Asp Ala
    1
   <210> 181
   <211> 47
   <212> DNA
   <213> Artificial Sequence
   <223> Description of Artificial Sequence: synthetic
         primer for generating mutant construct named
         MBPC125-SYEV
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| <400> 181 gacatetetg aagtgagtta ttaggeagaa tteegaeatg aeteagg   | 47 |
|---|----|
| <210> 182<br><211> 48<br><212> DNA<br><213> Artificial Sequence   |    |
| <220> <223> Description of Artificial Sequence: synthetic primer for generating mutant construct named MBPC125-SYEV |    |
| <400> 182 tgagtcatgt cggaattetg cetaataact caetteagag ateteete  | 48 |
| <210> 183 <211> 6 <212> PRT <213> Artificial Sequence   |    |
| <220> <223> Description of Artificial Sequence: synthetic peptide sequence  |    |
| <400> 183<br>Lys Lys Ser Tyr Glu Val<br>1 5   |    |
| <210> 184<br><211> 10<br><212> PRT<br><213> Artificial Sequence   |    |
| <220> <223> Description of Artificial Sequence: synthetic peptide sequence  |    |
| <400> 184<br>Val Glu Ala Asn Tyr Glu Val Glu Gly Glu<br>1 5   |    |
| <210> 185<br><211> 10<br><212> PRT<br><213> Artificial Sequence   |    |
| <220> <223> Description of Artificial Sequence: synthetic peptide sequence  |    |
| <400> 185<br>Val Glu Ala Asn Tyr Ala Val Glu Gly Glu<br>1 5   |    |
| <210> 186<br><211> 8<br><212> PRT<br><213> Artificial Sequence  |    |

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<223> Description of Artificial Sequence: synthetic
     peptide sequence
<400> 186
Asp Tyr Lys Asp Asp Asp Lys
<210> 187
<211> 4
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: synthetic
      peptide sequence
 <400> 187
 Ser Tyr Glu Ala
  1
 <210> 188
 <211> 4
 <212> PRT
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: synthetic
       peptide sequence
 <400> 188
 Ser Tyr Ala Val
   1
  <210> 189
  <211> 5
  <212> PRT
  <213> Artificial Sequence
  <223> Description of Artificial Sequence: synthetic
        peptide sequence
  <400> 189
  Val Ser Tyr Glu Ala
   <210> 190
   <211> 13
   <212> PRT
   <213> synthetic peptide sequence
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<400> 190
Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Trp Lys Lys
<210> 191
<211> 23
<212> PRT
<213> synthetic peptide sequence
 <400> 191
 Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu
 Val Glu Phe Arg Trp Lys Lys
 <210> 192
 <211> 15
 <212> PRT
 <213> synthetic peptide sequence
  <220>
  <221> SITE
  <222> (1)..(1)
  <223> amino acid at position 1 is biotinylated
   <220>
   <221> SITE
   <222> (14)..(14)
   <223> cys at position 14 is derivatized with an oregon green
   <400> 192
   Lys Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Lys Lys
    <210> 193
    <211> 22
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<212> PRT

<213> synthetic peptide sequence <220> <221> SITE <222> (1)..(1) <223> amino acid at position 1 is biotinylated <220> <221> SITE <222> (21)..(21) <223> cys at position 21 is derivatized with an oregon green <400> 193 Gly Leu Thr Asn Ile Lys Thr Glu Glu Ile Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Lys Lys <210> 194 <211> 6806 <212> DNA fusion protein comprising a maltose binding protein with 125 amino <213> acids from APP C-terminus. ccgacaccat cgaatggcgc aaaacctttc gcggtatggc atgatagcgc ccggaagaga 60 gtcaattcag ggtggtgaat gtgaaaccag taacgttata cgatgtcgca gagtatgccg 120 gtgtctctta tcagaccgtt tcccgcgtgg tgaaccaggc cagccacgtt tctgcgaaaa 180 cgcgggaaaa agtggaagcg gcgatggcgg agctgaatta cattcccaac cgcgtggcac 240 aacaactggc gggcaaacag tcgttgctga ttggcgttgc cacctccagt ctggccctgc 300 acgcgccgtc gcaaattgtc gcggcgatta aatctcgcgc cgatcaactg ggtgccagcg 360 tggtggtgtc gatggtagaa cgaagcggcg tcgaagcctg taaagcggcg gtgcacaatc 420 ttctcgcgca acgcgtcagt gggctgatca ttaactatcc gctggatgac caggatgcca 480 ttgctgtgga agctgcctgc actaatgttc cggcgttatt tcttgatgtc tctgaccaga 540

cacccatcaa cagtattatt ttctcccatg aagacggtac gcgactgggc gtggagcatc

tgccataacc atgagtgata acactgcggc caacttactt ctgacaacga tcggaggacc 4500 gaaggagcta accgettttt tgeacaacat gggggateat gtaactegee ttgategttg 4560 ggaaccggag ctgaatgaag ccataccaaa cgacgagcgt gacaccacga tgcctgtagc 4620 aatggcaaca acgttgcgca aactattaac tggcgaacta cttactctag cttcccggca 4680 acaattaata gactggatgg aggcggataa agttgcagga ccacttctgc gctcggccct 4740 teeggetgge tggtttattg etgataaate tggageeggt gagegtgggt etegeggtat 4800 cattgcagca ctggggccag atggtaagcc ctcccgtatc gtagttatct acacgacggg 4860 gagtcaggca actatggatg aacgaaatag acagatcgct gagataggtg cctcactgat 4920 taagcattgg taactgtcag accaagttta ctcatatata ctttagattg atttaaaact 4980 tcatttttaa tttaaaagga tctaggtgaa gatccttttt gataatctca tgaccaaaat 5040 cccttaacgt gagttttcgt tccactgagc gtcagacccc gtagaaaaga tcaaaggatc 5100 ttcttgagat cctttttttc tgcgcgtaat ctgctgcttg caaacaaaaa aaccaccgct 5160 accageggtg gtttgtttge eggateaaga getaecaaet etttteega aggtaaetgg 5220 cttcagcaga gcgcagatac caaatactgt ccttctagtg tagccgtagt taggccacca 5280 cttcaagaac tetgtageac egectacata eetegetetg etaateetgt taccagtgge 5340 tgctgccagt ggcgataagt cgtgtcttac cgggttggac tcaagacgat agttaccgga 5400 taaggcgcag cggtcgggct gaacgggggg ttcgtgcaca cagcccagct tggagcgaac 5460 gacctacacc gaactgagat acctacagcg tgagctatga gaaagcgcca cgcttcccga 5520 agggagaaag geggaeaggt ateeggtaag eggeagggte ggaaeaggag agegeaegag 5580 ggagcttcca gggggaaacg cctggtatct ttatagtcct gtcgggtttc gccacctctg 5640 acttgagcgt cgatttttgt gatgctcgtc aggggggggg agcctatgga aaaacgccag 5700 caacgeggee tttttaeggt teetggeett ttgetggeet tttgeteaca tgttetttee 5760 tgcgttatcc cctgattctg tggataaccg tattaccgcc tttgagtgag ctgataccgc 5820 tegeegeage egaaegaeeg agegeagega gteagtgage gaggaagegg aagagegeet 5880 gatgcggtat tttctcctta cgcatctgtg cggtatttca caccgcatat ggtgcactct 5940 cagtacaatc tgctctgatg ccgcatagtt aagccagtat acactccgct atcgctacgt 6000 gactgggtca tggctgcgcc ccgacacccg ccaacacccg ctgacgcgcc ctgacgggct 6060 tgtctgctcc cggcatccgc ttacagacaa gctgtgaccg tctccgggag ctgcatgtgt 6120 cagaggtttt caccgtcatc accgaaacgc gcgaggcagc tgcggtaaag ctcatcagcg 6180 tggtcgtgaa gcgattcaca gatgtctgcc tgttcatccg cgtccagctc gttgagtttc 6240 tccagaagcg ttaatgtctg gcttctgata aagcgggcca tgttaagggc ggttttttcc 6300 tgtttggtca cttgatgcct ccgtgtaagg gggaatttct gttcatgggg gtaatgatac 6360

| cgatgaaacg | agagaggatg | ctcacgatac | gggttactga | tgatgaacat | gcccggttac | 6420 |
|------------|------------|------------|------------|------------|------------|------|
| tggaacgttg | tgagggtaaa | caactggcgg | tatggatgcg | gcgggaccag | agaaaaatca | 6480 |
| ctcagggtca | atgccagcgc | ttcgttaata | cagatgtagg | tgttccacag | ggtagccagc | 6540 |
| agcatcctgc | gatgcagatc | cggaacataa | tggtgcaggg | cgctgacttc | cgcgtttcca | 6600 |
| gactttacga | aacacggaaa | ccgaagacca | ttcatgttgt | tgctcaggtc | gcagacgttt | 6660 |
| tgcagcagca | gtcgcttcac | gttcgctcgc | gtatcggtga | ttcattctgc | taaccagtaa | 6720 |
| ggcaaccccg | ccagcctagc | cgggtcctca | acgacaggag | cacgatcatg | cgcacccgtg | 6780 |
| gccaggaccc | aacgctgccc | gaaatt     |            |            |            | 6806 |
|            |            |            |            |            |            |      |

<210> 195

<211> 13

<212> PRT

<213> synthetic peptide sequence

<220>

<221> MOD\_RES

<222> (1)..(1)

<223> ACETYLATION (MCA)

<220>

<221> SITE

<222> (11)..(11)

<223> 2,4-dinitrophenyl group after the Lys at position 11

<400> 195

Ser Glu Val Asn Leu Asp Ala Glu Phe Arg Lys Arg Arg 1 5 10

<210> 196

<211> 12

<212> PRT

<213> synthetic peptide sequence

<220>

<221> SITE

<222> (4)..(4)

<223> amino acid at position 4 has been derivatized with a statine

<400> 196

<210> 197

<211> 10

<212> PRT

<213> synthetic peptide sequence

<220>

<221> SITE

<222> (4)..(4)

<223> amino acid at position 4 has been derivatized with a statine

<220>

<221> SITE

<222> (10)..(10)

<223> amino acid at position 10 has been derivatized with Bodipy FL

<400> 197

Ser Glu Val Asn Val Ala Glu Phe Arg Cys 1 5 10